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Sequence Listing could not be accepted.

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Reviewer: Durreshwar Anjum

Timestamp: [year=2008; month=10; day=29; hr=14; min=12; sec=15; ms=896;
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Reviewer Comments:

<110> APPLICANT: Hakoto KOBAYAHSI

Yugo HABATA

Ryo FUJII

Shuji HINUMA

<120> TITLE OF INVENTION: Methods of Screening for Ligands for FPRL2

<130> FILE REFERENCE: 3171 US0P

<140> CURRENT APPLICATION NUMBER:10554234

<141> CURRENT FILING DATE:2005-10-21

<150> PRIOR APPLICATION NUMBER: PCT/JP2004/005829

<151> PRIOR FILING DATE: 2004-04-22

<150> PRIOR APPLICATION NUMBER: JP 2003-118760

<151> PRIOR FILING DATE: 2003-04-23

<160> NUMBER OF SEQ ID NOS: 8

<210> SEQ ID NO 1

<211> LENGTH: 353

<212> TYPE: PRT

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 1

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Application No: 10554234 Version No: 2.0

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Output Set:

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Finished: 2008-09-30 12:20:14.257
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Total Errors: 0
No. of SeqIDs Defined: 8
Actual SeqID Count: 8

Error code	Error Description
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W 402	Undefined organism found in <213> in SEQ ID (4)
W 333	tabs used in amino acid numbering SEQID (8)

<110> APPLICANT: Hakoto KOBAYAHSI

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<210> SEQ ID NO 1

<211> LENGTH: 353

<212> TYPE: PRT

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 1

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Pro Glu Pro Ala Gly His Thr Val Leu Trp Ile Phe Ser Leu Leu Val
      20              25              30
His Gly Val Thr Phe Val Phe Gly Val Leu Gly Asn Gly Leu Val Ile
      35              40              45
Trp Val Ala Gly Phe Arg Met Thr Arg Thr Val Asn Thr Ile Cys Tyr
      50              55              60
Leu Asn Leu Ala Leu Ala Asp Phe Ser Phe Ser Ala Ile Leu Pro Phe
      65              70              75              80
Arg Met Val Ser Val Ala Met Arg Glu Lys Trp Pro Phe Ala Ser Phe
      85              90              95
Leu Cys Lys Leu Val His Val Met Ile Asp Ile Asn Leu Phe Val Ser
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Val Tyr Leu Ile Thr Ile Ile Ala Leu Asp Arg Cys Ile Cys Val Leu
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His Pro Ala Trp Ala Gln Asn His Arg Thr Met Ser Leu Ala Lys Arg
      130             135             140
Val Met Thr Gly Leu Trp Ile Phe Thr Ile Val Leu Thr Leu Pro Asn
      145             150             155             160
Phe Ile Phe Trp Thr Thr Ile Ser Thr Thr Asn Gly Asp Thr Tyr Cys
      165             170             175
Ile Phe Asn Phe Ala Phe Trp Gly Asp Thr Ala Val Glu Arg Leu Asn
      180             185             190
Val Phe Ile Thr Met Ala Lys Val Phe Leu Ile Leu His Phe Ile Ile
      195             200             205
Gly Phe Thr Val Pro Met Ser Ile Ile Thr Val Cys Tyr Gly Ile Ile
      210             215             220
Ala Ala Lys Ile His Arg Asn His Met Ile Lys Ser Ser Arg Pro Leu
      225             230             235             240
Arg Val Phe Ala Ala Val Val Ala Ser Phe Phe Ile Cys Trp Phe Pro
      245             250             255
Tyr Glu Leu Ile Gly Ile Leu Met Ala Val Trp Leu Lys Glu Met Leu
      260             265             270
Leu Asn Gly Lys Tyr Lys Ile Ile Leu Val Leu Ile Asn Pro Thr Ser
      275             280             285
Ser Leu Ala Phe Phe Asn Ser Cys Leu Asn Pro Ile Leu Tyr Val Phe
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290	295	300
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305	310	315
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	325	330
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		350
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<210> SEQ ID NO 2

<211> LENGTH: 1059

<212> TYPE: DNA

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 2

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gcattctggg	gtgacactgc	tgtagagagg	ttgaacgtgt	tcattaccat	ggccaaggtc	600
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ctctacgtct	ttatgggtcg	taacttccaa	gaaagactga	ttcgctcttt	gcccactagt	960
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<210> SEQ ID NO 3

<211> LENGTH: 6

<212> TYPE: PRT

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: amino acid sequence of GHRP-6

<220> FEATURE:

<223> OTHER INFORMATION: Trp is a D-form

<400> SEQUENCE: 3

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<210> SEQ ID NO 4

<211> LENGTH: 11

<212> TYPE: PRT

<213> ORGANISM: Aplysia sp.

<400> SEQUENCE: 4

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<210> SEQ ID NO 5

<211> LENGTH: 12

<212> TYPE: PRT
<213> ORGANISM: Sus scrofa
<400> SEQUENCE: 5
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<210> SEQ ID NO 6
<211> LENGTH: 36
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 6
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Leu Ala Arg Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr
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Arg Gln Arg Tyr
35

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<211> LENGTH: 10
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 7
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<210> SEQ ID NO 8
<211> LENGTH: 6
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 8
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1 5